

FIG. 1A

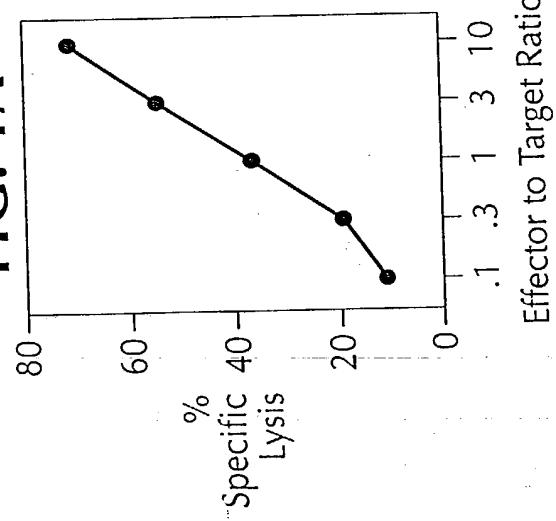


FIG. 1B

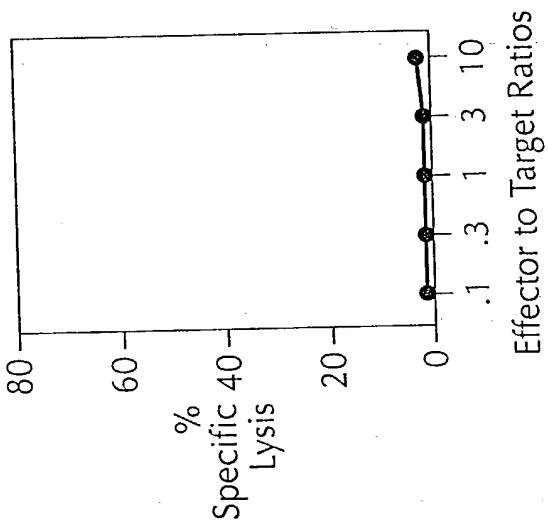


FIG. 1C

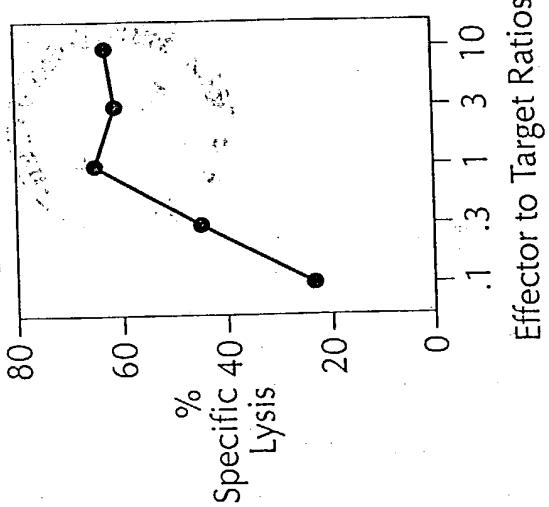


FIG. 1D

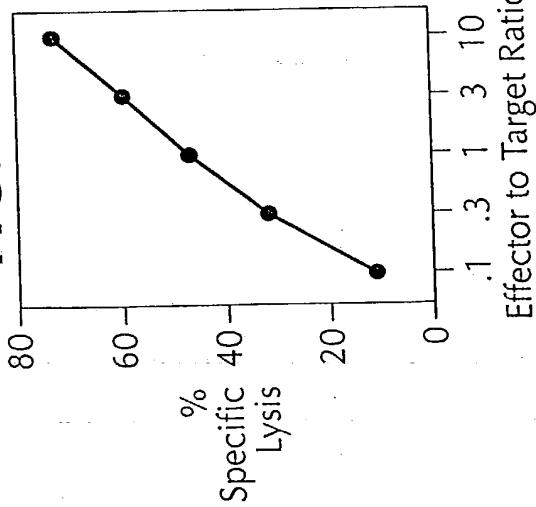


FIG. 1E

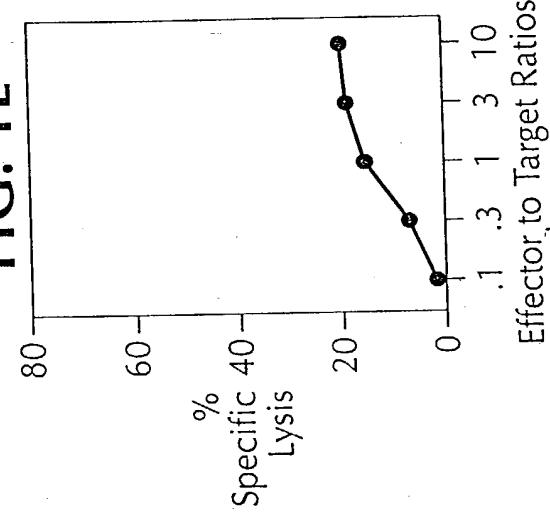


FIG. 1F

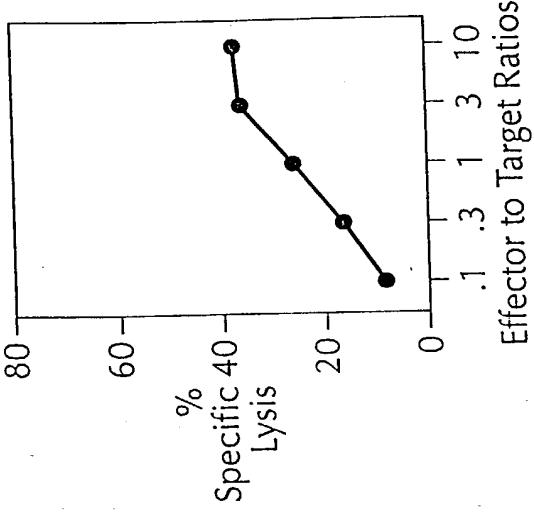


FIG. 2

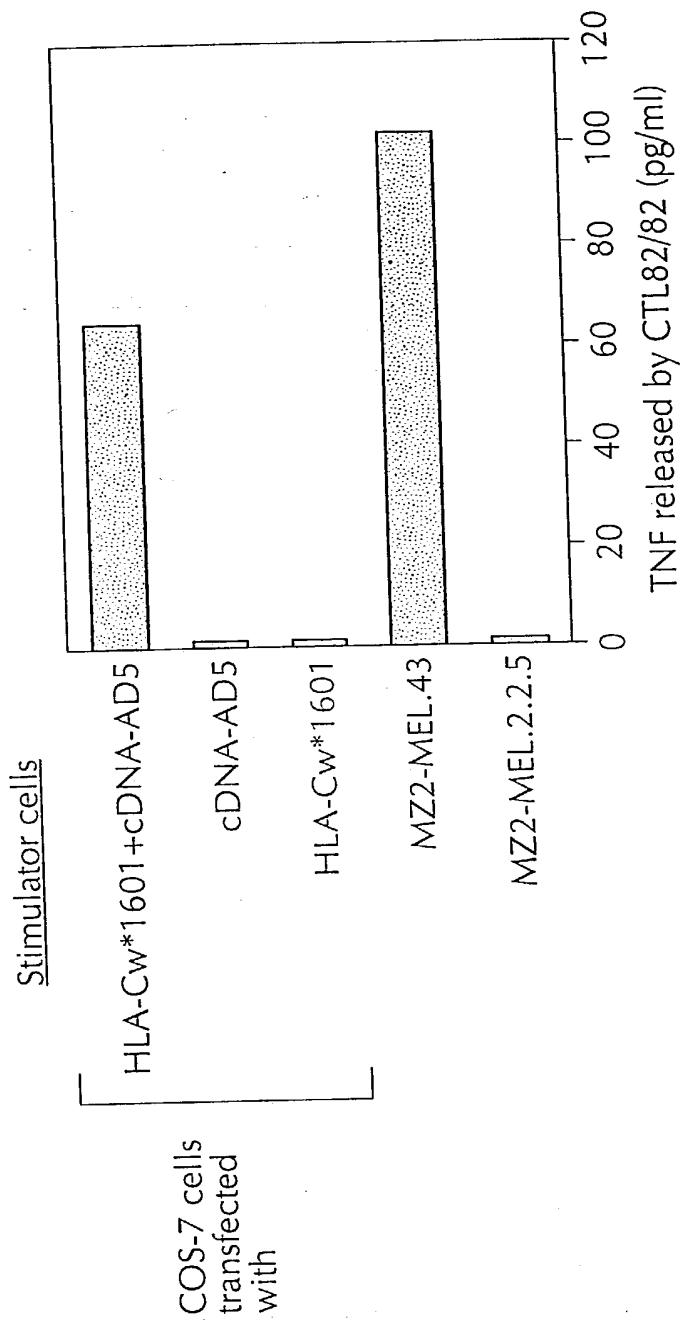


FIG. 3A

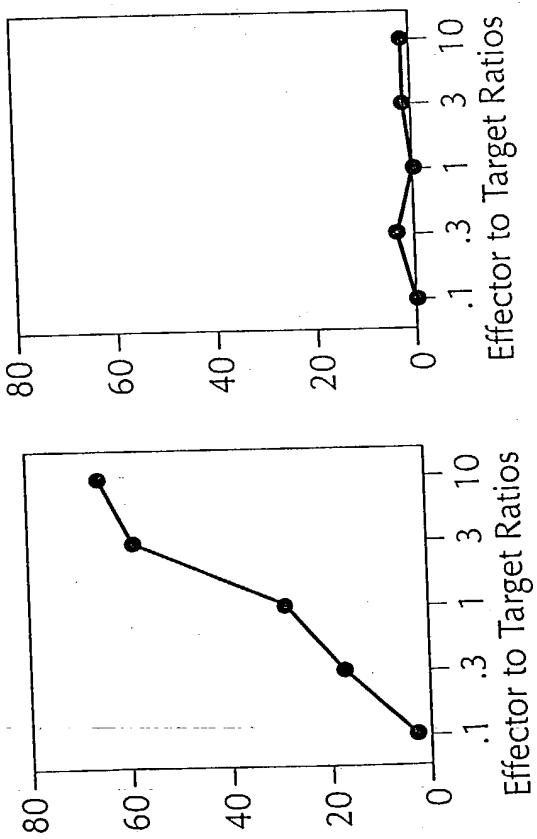


FIG. 3B

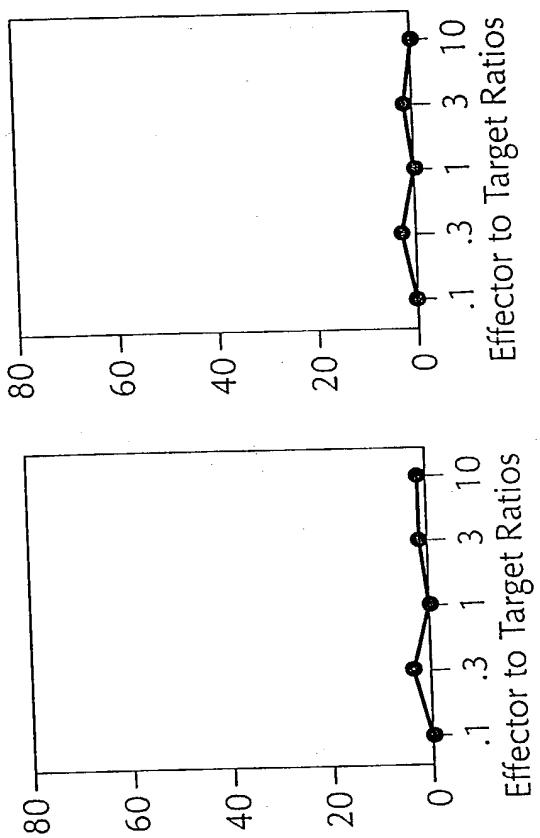


FIG. 3C

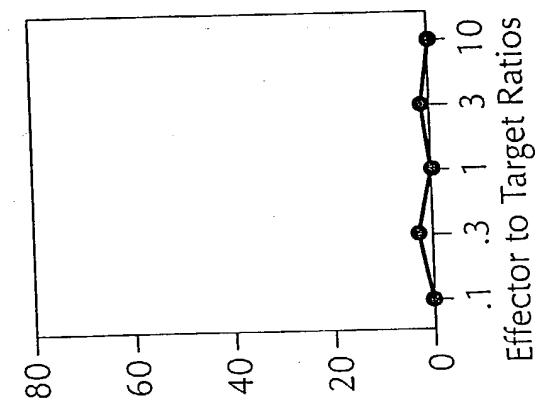


FIG. 3D

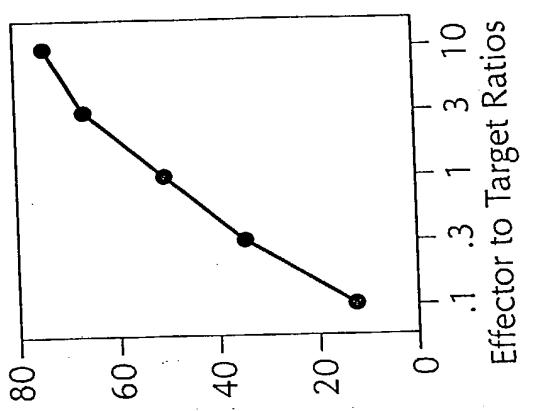


FIG. 4

CGCCAATTAA GGGTCTCCGG TATCTCCCGC TGAGCTGCTC TGTTCGGGC TTAGAGGACCC 60
 AGGAGAAGGG GGAGCTGGAG GCTGGAGCCCT GTAACACCCGT GGCTCGTCTC ACTCTGGATG 120
 GTGGTGGCAA CAGAGATGGC AGCCGAGCTG GAGTGTAGG AGGGCGGGCCT GAGCGGTAGG 180
 M A A R A V F L A L S A Q S 13
 AGTGGGGCTG GAGCAGTAAG ATGGCGGCCA GAGCGGTTT TCTGGCATTC TCTGGCCAGC 240
 L L Q A R L M K E E S P V V S W R L E P 33
 TGCTCCAAGC CAGGCTGATG AAGGAGGAGT CCCCTGTGGT GAGCTGGAGG TTGGAGCCTG 300
 E D G T A L C F I F 43
 AAGACGGCAC AGCTCTGTGC TTCATCTTCT GAGGTTGTGG CAGCCACGGT GATGGAGACG 360
 GCAGCTAAC AGGAGCAATA GGAGGAGATG GAGTTCACT GTGTCAGCCA GGATGGTCTC 420
 GATCTCCTGA CCTCGTGA CGCCCGCCTT GGCCCTTCCAA AGTGCCGAGA TTACAGCGAT 480
 GTGCATTTCG TAAGCACTT GGAGCCACTA TCAAAATGCTG TGAAAGAGAAA TGTACACCCAGA 540
 TGTATCATT TAAGCACTT GCAGGAGCCG GCTCCTTCA GGATTTCAGT CACATCTTC 600
 TGCTTGTCC AGAACACATT GACCAAGCTC CTGAAAGATG TAAGTTTACT ACGCATAGAC 660
 TTTTAAACTT CAACCAATGT ATTACTGAA AATAACAAAT GTTGTAAATT CCCTGAGTGT 730
 TATTCTACTT GTATTAAGA GTAAATAATAC ATAATCATTAA AAATCTGAGG GATCATGGCC 780
 AGAGATGTGTT GGGGAGGAA ATGTTATCAA CGGTTTCATT GAAATTAAT CCAAAAAGTT 840
 ATTTCCTCAG AAAATCAA TAAAGTTTGC ATGTTTTTTA TTCTTAAAC ATTAAATAA 900
 CCACTGTAGA ATGATGTAAA TAGGGACTGT GCAGTATTTC TGACATATAC TATAAAATTA 960
 TTAAAAAGTC AATCAGTATT CAACATCTT TACACTAAA AGCC 1004

FIG. 5

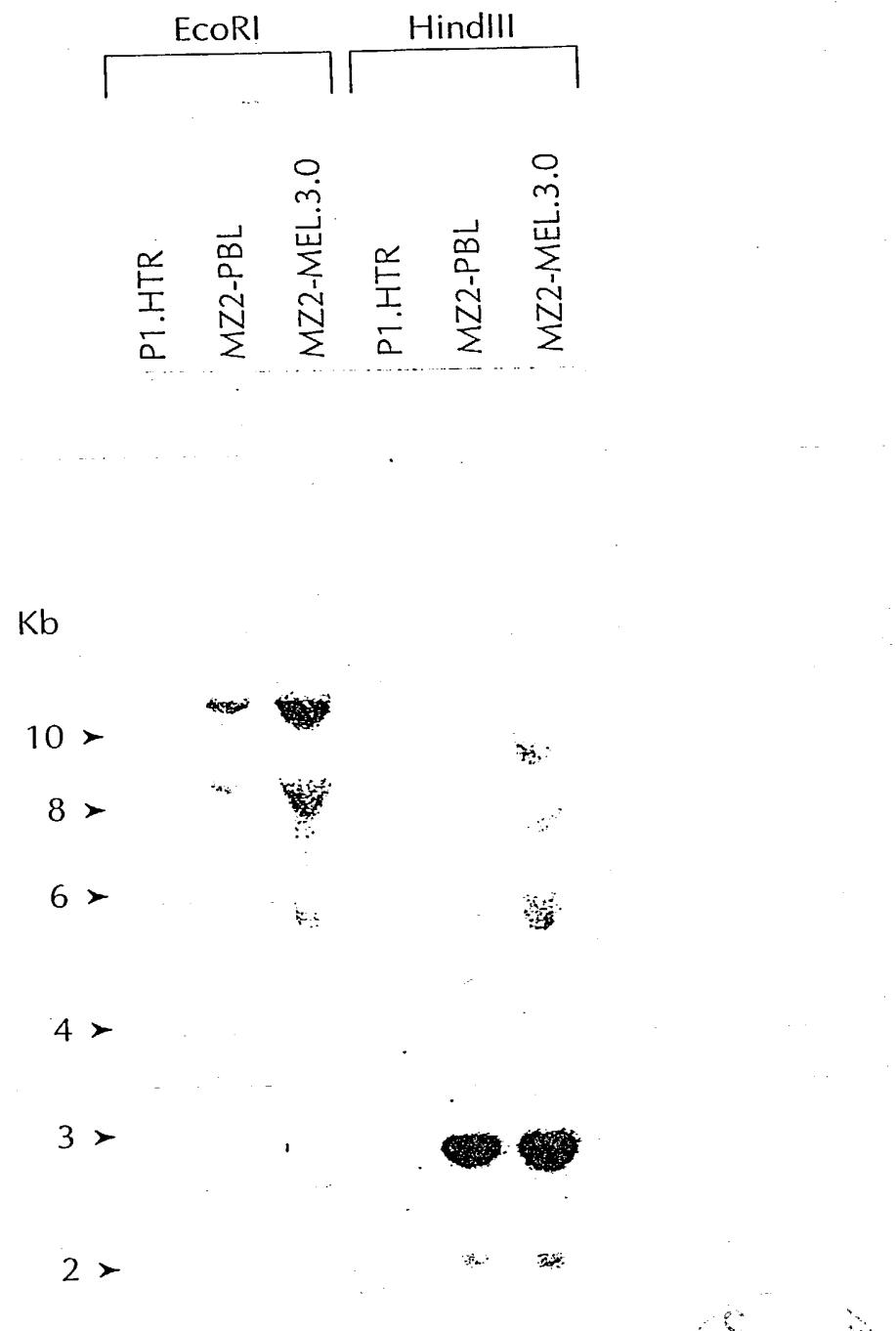


FIG. 6

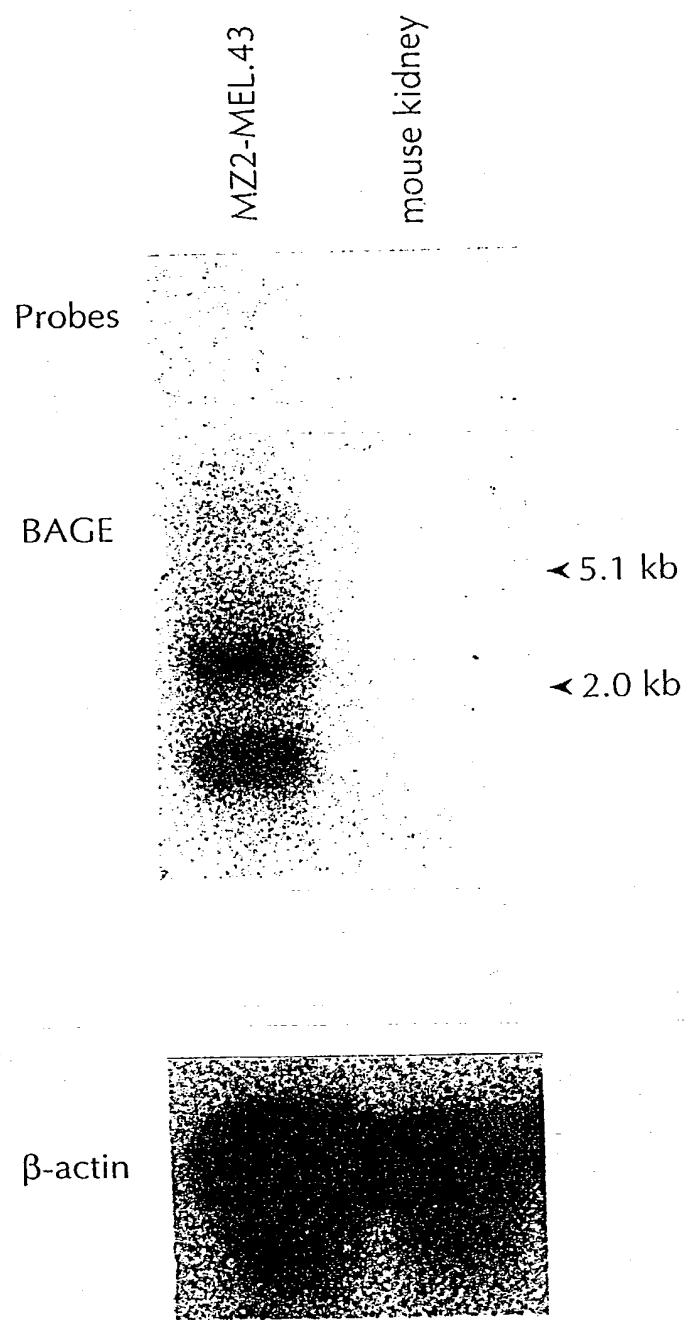


FIG. 7

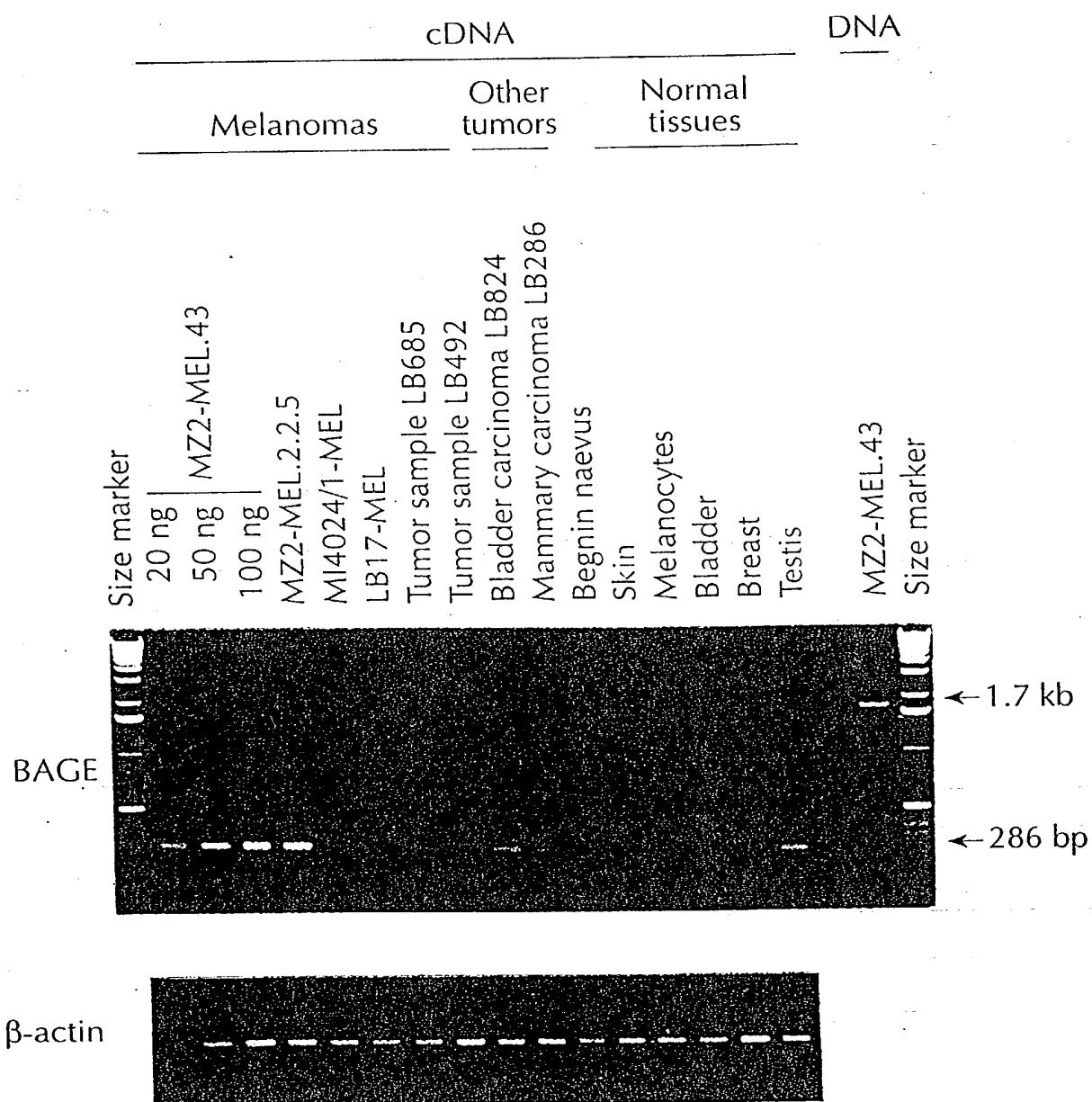


FIG. 8

